09/881569

FILE 'REGISTRY' ENTERED AT 14:29:32 ON 15 NOV 2002 L1 0 S BTS.PCGAVMILYFHKWR/SQSP

L2 0 S PCGAVMILY/SQSP

FILE 'HOME' ENTERED AT 14:32:35 ON 15 NOV 2002

Searcher: Shears 308-4994

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| 990S-0127462. 990S-0128234. 990S-0128714. 990S-0129845. 990S-0130077. | 99US-0125788. 99US-0126264. 99US-0126785. | 99US-0121825. 99US-0123180. 99US-0123548. | EP-0301439. | | | ana. | genetic mapping; gene expression control; | Protein identification; signal transduction pathway; metabolic pathway; | Arabidopsis thaliana protein fragment SEQ ID NO: 3815. | (first entry) | | ; Protein; 281 AA. |

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                   The present invention describes substantially purified human proteins (referred to as open reading frame, ORFX, where X is 1-11491 (see Table 1 in the specification). ABN15762 to ABN27252 encode the human ORFX proteins given in ABP00010 to ABP11500. ORFX proteins are useful for treating or preventing a pathology associated with an ORFX-associated disorder in humans, and in the manufacture of a medicament for treating a syndrome associated with ORFX-associated disorder. ORFX polynucleotide sequences can be used in gene therapy. ORFX sequences can be used in the treatment of cancer, hyperproliferative disorders, cirrhosis of liver, psoriasis, benign tumours, keloid, degenerative disorders, haemorrhage,
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                                                                                                                   (WISC ) WISCONSIN ALUMNI RES (PION-) PIONEER HI-BRED INT 1 (MINU ) UNIV MINNESOTA.
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1.8;
                                                                           TG,
                                                                           Phillips
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RESULT 6
AAU78464
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               용
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CC The invention relates to plant genetic engineering. More specifically, CC the invention relates to isolated and purified polycomb group (PCC) genes CC from Zea mays (maize), termed Mez1 and Mez2 (Maize E(2)-like 1 and 2) CC genes comprising a sequence of 3180 and 3030 bp respectively, defined in CC the specification, or its conservatively modified and polymorphic CC variants. The PCG genes of the invention are useful in constructing CC recombinant expression cassettes which can be introduced into a desired CC plant, to produce a transgenic plant. Promoters derived from the Mez1 or CC Mez2 genes are useful to direct expression of heterologous sequences. The CC modified variants of the PCG genes are useful for generating or selecting CC antibodies immunoreactive to the non-variant polypeptide and to follow the segregation of chromosome regions and in marker assisted selection CC methods for crop improvement. Mez1 and Mez2 proteins can be used to CC repress the expression or prevent the expression of a desired target gene expression in plants to create novel plants having improved agronomic traits. Protein derived from a PCG gene of the invention is useful as an immunogen for producing monoclonal or polyclonal antibodies which are CC useful for screening plants for the expression of a Mez1 or Mez2 CC polypeptide, and for affinity chromatography for isolating a Mez1 or Mez2 CC invention. This sequence represents the protein of Mez 2 of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
New polypeptide with sphingolipid ceramide
                                 N-PSDB;
                                                                                    Ito
                                                                                                                  (TAKI )
                                                                                                                                                   26-SEP-2000;
                                                                                                                                                                                   26-SEP-2001;
                                                                                                                                                                                                                      04-APR-2002
                                                                                                                                                                                                                                                                                       Shewanella alga
                                                                                                                                                                                                                                                                                                                                   Sphingolipid ceramide deacylase; nootropic; neuroprotective; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                            02-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU78464 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Novel polycomb group nucleic acids and polypeptides from maize useful for producing transgenic plants and to repress expression or prevent the expression of a desired target gene in specific tissue in plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 317
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 BTSZPCGAVMILYFHKWR 18
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                                                 2002-340011/37
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7; Conserv
                                 ABK47644
                                                                                                                  TAKARA SHUZO CO LTD
                                                                                Furusato M,
                                                                                                                                                                                                                                                                                                                    neurodegenerative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                   2000JP-0293181
                                                                                                                                                                                                                                                                                                                                                                          ceramide
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                                                                                Sueyoshi
                                                                                                                                                                                                                                                                                                                                                                       deacylase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           954 AA
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Pred. No.
                                                                                                                                                                                                                                                                                                                       disease;
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                                                                                                                                                                                                                                                                                                                     leukaemia; wound; enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95;
deacylase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 893;
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activity
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RESULT 7
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                        The invention relates to a polypeptide with sphingolipid ceramide deacylase activity and its encoding nucleic acid. The polypeptide and its encoded gene are applicable in studying structure and function of sphingolipid and its engineering for treatment of e.g. neurodegenerative diseases, leukaemia and wounds. The present sequence represents the amino acid sequence of sphingolipid ceramide deacylase #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a polypeptide with sphingolipid ceramide deacylase activity and its encoding nucleic acid. The polypeptide and its encoded gene are applicable in studying structure and function of sphingolipid and its engineering for treatment of e.g. neurodegenerative diseases, leukaemia and wounds. The present sequence represents the amino acid sequence of sphingolipid ceramide deacylase.
                                                                                                                                                        New polypeptide with sphingolipid ceramide deacylase activity and encoded gene, applicable in studying structure and function of sphingolipid and its engineering for treatment of e.g. neurodegenerative diseases and leukaemia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           encoded gene, applicable in studying structure and function sphingolipid and its engineering for treatment of e.g. neurodegenerative diseases and leukaemia -
Sequence
                                                                                                                                                                                                                                                                          Ito
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Shewanella alga.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sphingolipid ceramide deacylase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 6; Page 59-64; 89pp; Japanese.
                                                                                                                              Example 2;
                                                                                                                                                                                                                                  N-PSDB;
                                                                                                                                                                                                                                                                                                    (TAKI ) TAKARA SHUZO CO
                                                                                                                                                                                                                                                                                                                                  26-SEP-2000; 2000JP-0293181
                                                                                                                                                                                                                                                                                                                                                             26-SEP-2001;
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                                                                                                                                                                                                                                                                                                                                                                                                                      WO200226963-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              vulnerary;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sphingolipid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAU78470 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     929
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2 TSZPCGAVMILYFH-----KWR 18
                                                                                                                                                                                                                                              2002-340011/37.
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                                                                                                                                                                                                                                  ABK47651.
                                                                                                                                                                                                                                                                         Furusato M,
                                                                                                                           Page 76-78; 89pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              neurodegenerative disease;
 992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      954 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ceramide deacylase
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         42.2%;
                                                                                                                                                                                                                                                                          Sueyoshi N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              992 AA
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          #2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nootropic; neuroprotective; cytostatic;
ease; leukaemia; wound; enzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 2-
1.5e+02;
5;
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Query Match Best Local S Matches 8

Similarity 8; Conser

Conservative

42.28;

Score 43; DB Pred. No. 1.6e 4; Mismatches

DB 23; 1.6e+02; 5;

Length 992;

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TSZPCGAVMILYFH----KWR

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RESULT 8
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                                                             Matches
                                                                          Query Match
Best Local
                                                                                                                                                               Mucociliary dysfunction is the inability of ciliated epithelium to clear mucus and sputum in lung airways. Mucociliary dysfunction is a serious complication of chronic obstructive lung diseases such as Chronic Bronchitis (CB), Bronchiectasis (BE), asthma and Cystic Fibrosis (CF). In addition, patients suffering from mucociliary dysfunction are susceptible to secondary bacterial infections. The protein sequence of human placental bikunin has been isolated (see AAB14159). Placental bikunin is a Kunitz-type serine protease inhibitor protein, which can stimulate the rate of mucociliary clearance of mucus and sputum in lung airways. Therefore, placental bikunin may be used for treating lung diseases such as CF, CB, BE, and chronic sinusitis and shue ear which are caused by retention and accumulation of mucus. The present sequence is a known Kunitz-like domain and was used to identify the known Kunitz-like domain and was used to identify the
                                                                                                                                                                                                                                                                                                                                                                                                                                            Stimulating mucociliary clearance rate of mucus and sputum in lung airways for treating lung diseases such as cystic fibrosis and bronchitis involves administering a Kunitz-type serine protease inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 22; 173pp; English
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                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2000-452127/39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hall
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-DEC-1998;
17-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BE; asthma; cystic fibrosis; CF; bacterial infection; placental bikunin; Kunitz-type serine protesse inhibitor; chronic sinusitis; glue ear; tissue factor pathway inhibitor precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mucociliary dysfunction; mucus; sputum;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tissue factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-FEB-2001
                                                                                                                                                     Kunitz-domains of placental bikunin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (FARB ) BAYER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-DEC-1999;
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12
                                                           Local Similarity
nes 5; Conserv
                              G
PCRALLLRYYYRYR 25
                           PCGAVMILYFHKWR 18
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                                                                                                                        57
                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CT,
                                                                                                                        AA;
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99US-0441966.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Newton BB,
                                                                        41.2%;
35.7%;
                                                        Score 42; DB
Pred. No. 11;
7; Mismatches
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAY34584-Y35879 represent the proteins encoded by all the open reading frames in the complete genome (see AAX91990) of Chlamydia pneumoniae. C. pneumoniae causes respiratory disease such as pneumonia and bronchitis and is thought to be a contributing factor in heart disease, sarcoidosis, sinusitis, purulent otitis media, erythema nodosum or pharyngitis. The polypeptides encoded by the open reading frames of the C. pneumoniae genome (see AAY34584-Y35879) can be used in immunogenic compositions as vaccines. Vectors containing C. pneumoniae nucleotides sequences can also be used as immunogenic compositions, especially where the vector directs the expression of a neutralising epitope of C. pneumoniae.
                                                                             Protease inhibitor; antitrypsin activity; shock therapy; kallikrein inhibitor; elastase inhibitor; arthritis ther
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-357842/30.
                                                                                                                                                                Sequence
                                                                                                                                                                                                                         28-NOV-1991
                                                                                                                                                                                                                                                                                AAP50700;
                                                                                                                                                                                                                                                                                                                                   AAP50700 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Griffais R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  04-NOV-1998;
21-NOV-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                vaccine; neutralising epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sinusitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Respiratory disease; pneumonia; bronchitis; heart disease; sarcoidosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Amino acid sequence of a Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-SEP-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (GEST ) GENSET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-NOV-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 TSZPCGAVMILYFHKWR 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTEPCSSIHISLNNDWR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1278-1279; Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity
6; Conser
                                                                                                                                                                of Gly(15)-Aprotinin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                purulent otitis media; erythema
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98US-0107078
97FR-0014673
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98WO-1B01890
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                41.2%;
                                                                                                                                                                                                                                                                                                                                      58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1912pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 42; DB Pred. No. 48; 5; Mismatches
                                                                                                                                                                                                                                                                                                                                      ₽
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ζ
                                                                             arthritis therapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nodosum; pharyngitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      6,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 223;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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Sus scrofa domestica

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Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                             AAW35008
                                                                                                                                                                                                                                                                                                                                       RESULT 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               replaced by one of (all I forms where applicable) Gly.Ala,Val,Leu, Ile,Met,Arg, alpha-aminobutyric acid, norvaline, Nle, dehydroalanine or homoserine. Also new are the intermediate derive. in which (1) 15-Lys is replaced by one of the specified Aas or their alkyl esters and any additional COOH in side chains or terminal posn. are partly or completely esterified; (2) the bond between 15-Lys and 16-Ala is open and only 15-Lys COOH is in free form, all other side chain COOH (3- and 50-Asp; 7- and 49-Glu) and C-terminal COOH (58-Ala) are present as alkyl esters; (3) the 15/16 bond is broken; 15-Lys is absent, the side chain and terminal COOH are esterified and only 14-Cys has a COOH in free form.
                                                                                                                                                                                Endoglucanase; cellulase; carboxymethylcellulose; cellulose; biomass; beta-1,4-glycosidic bond; hydrolysis; saccharification; thermostable enzyme; thermophilic; glycosidase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 1; 51pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-NOV-1983;
28-JUL-1983;
                                                                                                                                                            Vibrio harveyi
                                                                                                                                                                                                                                                             21-MAY-1998
                                                                                                                                                                                                                                                                                                              AAW35008 standard; Protein; 504 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New aprotinin derivs. with 15-lysine new intermediate, useful as protease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1985-039382/07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tschesche H, Wenzel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (FARB ) BAYER AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-FEB-1985
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EP132732-A.
                                   (RECO-) RECOMBINANT BIOCATALYSIS INC
                                                           22-MAY-1996;
                                                                                    22-MAY-1997;
                                                                                                             27-NOV-1997
                                                                                                                                     WO9744361-A1
                                                                                                                                                                                                                                    Vibrio harveyi endoglucanase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The inventors claim homologues of aprotinin in which 15-Lys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         treating shock
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-JUL-1984;
                                                                                                                                                                                                                                                                                                                                                                            13
                                                                                                                                                                                                                                                                                                                                                                                                  5 PCGAVMILYFH 15
                                                                                                                                                                                                                                                                                                                                                                          PCGARIIRYFY 23
             Mathur EJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            58
                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Α,
                                                                                                                                                             (Clone 91GP2)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  83DE-3339693
83DE-3327277
                                                           96US-0651572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        84EP-0108293
                                                                                    97WO-US08793
                                                                                                                                                                                                                                                                                                                                                                                                                                      40.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ξ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Schmuck
                                                                                                                                                                                                                                                                                                                                                                                                                          Score 41; DB
Pred. No. 17;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                            <u>ب</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Schnabel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     replaced by inhibitors,
                                                                                                                                                                                                                                                                                                                                                                                                                                         DB
17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  6,
                                                                                                                                                                                                                                                                                                                                                                                                                          2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 58
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e.g. for
                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         has been
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                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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RESULT 12
AAW59445
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
                                                                                                                                                                                                           07-MAR-1997;
29-JUL-1996;
30-OCT-1996;
                                                                       production resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        host cells, especially prokaryotic host cells transformed with a plasmid or virus derived vector including the endoglucanase DNA (see AAT94216). 24 Endoglucanases (see AAW34986-W35008) are claimed. They can be used to degrade cellulose for the conversion of plant biomass into fuels and chemicals, for use in detergents, textiles, animal feed, waste treatment, and in the fruit juice and brewing industries for the clarification and extraction of juices.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (Clone 91GP2) that is capable of degrading carboxymethylcellulose and of hydrolysing the beta-1,4-glycosidic bonds in cellulose. I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; Fig 1x; 164pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Endoglucanase(s), preferably form archael bacterium, AEPII la
useful to degrade carboxymethylcellulose and hydrolyse of
                                                                                                                        N-PSDB;
                                                                                                                                                            Bueschges R,
                                                                                                                                                                                                                                                                29-JUL-1997;
                                                                                                                                                                                                                                                                                         05-FEB-1998
                                                                                                                                                                                                                                                                                                                                                                MLO; mildew; pathogen; resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW59445;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW59445 standard; Protein; 544 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                has homology to an endoglucanase of archaebacterium AEPIIIa (see AAW34985). It can be produced from native cells or from recombin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  beta-1,4-glycosidic bonds
                                            Claim 8; Fig 14; 150pp; English
                                                                                                New isolated
                                                                                                                                    WPI; 1998-159149/14.
                                                                                                                                                                                     (INNE-) INNES
                                                                                                                                                                                                                                                                                                                                       Hordeum vulgare
                                                                                                                                                                                                                                                                                                                                                                                         Hordeum vulgare MLO protein homologue
                                                                                                                                                                                                                                                                                                                                                                                                                 13-OCT-1998 (first entry)
                                                                                                                                                                                                                                                                                                               WO9804586-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
mes 6; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 PCGAVMILYFHKW 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PDGGVVVGYWHNW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein comprises an endoglucanase of Vibrio harveyi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1998-018435/02.
                                                                                                                        AAV35030.
                                                                                 ed Mlo gene of of transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   504 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                            Panstruga R,
                                                                                                                                                                                     CENT INNOVATIONS
                                                                                                                                                                                                             97GB-0004789.
96GB-0015879.
96GB-0022626.
                                                                                                                                                                                                                                                               97WO-GB02046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             40.2%;
46.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  46
                                                                                 barley
plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    in cellulose
                                                                                                                                                             Schulzelefert PMJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ω
•
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 41;
Pred. No.
                                                                                                                                                                                     LTD JOHN

    used to develop products for the which have increased pathogen

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 19;
. 1.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 504;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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The sequence is that encoded by a homologue of the MLO gene, wild-type Mlo exerts a negative regulatory function on a pathogen defence response, such that mutants exhibit a defence response in the absence of pathogen.

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RESULT 13
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            the treatment, prevention and diagnosis of medical conditions caused by P. acnes. The disorders include SAPHO syndrome (synovitis, acne, pustulosis, hypertosis and ostcomyelitis), uveitis and endophthalmitis. P. acnes is also involved in infections of bone, joints and the central nervous system, however it is particularly involved in the inflammatory lesions associated with acne vulgaris. A method for detecting the presence or absence of P. acnes in a patient comprises contacting a sample with a binding agent that binds to the proteins of the invention and determining the amount of bound protein in the sample. The polypeptides may be used as antigens in the production of antibodies specific for P. acnes proteins. These antibodies can be used to
                                                                                                                                                                                                                                             Sequences AAU39105-AAU68017 represent Propionibacterium acnes polypeptides. The proteins and their associated DNA sequences
                                                                                                                                                                                                                                                                                                                                                 Propionibacterium acnes polypeptides and nucleic acids useful for vaccinating against and diagnosing infections, especially useful treating acne vulgaris -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   21-APR-2000; 2000US-199047P.
02-JUN-2000; 2000US-208841P.
07-JUL-2000; 2000US-216747P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       stimulate a defence response in transgenic plants conferring pathogen resistance, especially resistance to powdery mildew The product can also be used for identifying compounds able t stimulate a defence response in a plant by interaction with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Down-regulation or out-competition of Mio function may
stimulate a defence response in transgenic plants confe
pathogen resistance, especially resistance to powdery m
The product can also be used for identifying compounds
                                                                                                                                                                                                                            polypeptides. The proteins and the treatment, prevention and d
                                                                                                                                                                                                                                                                                                             Example 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                       N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   L'maisonneuve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Skeiky YAW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WO200181581-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Propionibacterium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     dermatological;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SAPHO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Propionibacterium acnes immunogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  27-FEB-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      encoded polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-APR-2001; 2001WO-US12865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAU42094;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAU42094 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CORI-) CORIXA CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        vveitis; endophthalmitis;
inflammatory lesion; acne
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8 AVMIL-----YFHKWR 18
                                                                                                                                                                                                                                                                                                                                                                                                                                     2001-616774/71.
DB; AAS59516.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AVMILVSVAMEHALHKLGHWFHKWR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               syndrome; synovitis; acne;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                               SEQ ID No
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  544 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Persing DH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
  expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ŗ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   osteopathic; neuroprotectant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zhang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  40.2%;
                                                                                                                                                                                                                                                                                                         3289; 1069pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mitcham JL, Wang SS,
, Jen S, Carter D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      bone; joint; central nervous system; vulgaris; enzyme linked immunosorbent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 41;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               pustulosis; hypertosis; osteomyelitis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       protein #2990
o.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 19;
1.8e+02;
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               The present invention relates to the genome sequence of Listeria monocytogenes EGD-e (see ABA03041). The genome sequence and fragments of it are useful for selecting probes and primers for detecting genes in L. monocytogenes and related organisms, and for studying genetic polymorphisms and other genomes. The present sequence is a protein encoded by the genome sequence of the present invention. Proteins expressed from the genome sequence are useful for raising specific antibodies, identification of L. monocytogenes and related organisms, and for biosynthesis and biodegradation, especially biosynthesis of Vitamin B12. The genome sequence and proteins encoded by the genome sequence and proteins encoded by the realso useful for selecting compounds that regulate gene expression and cell replication and modulate L. monocytogenes related diseases. In addition, the genome
                                                                                                                                                                                                                                                                                                                                                                                                      Genomic sequence for Listeria monocytogenes, useful and prevention of Listeria and related bacterial inf related polypeptides -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI;
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Note: The sequence data for this patent did no specification, but was obtained in electronic
                                                                                                                                                                                                                                                                                                                                                          Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antibacterial; gene therapy; vaccine; biosynthesis; biodegradation; vitamin Bl2; bacterial infection; disease.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Rose M,
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                                         This invention describes novel transgenic mice expressing proteins related to the pathology of Alzheimer's disease and which provide models for studying potentially therapeutic compounds. The transgenic mice contain a DNA sequence encoding a beta-amyloid precursor protein (APP) and a nerve tissue specific promoter operably linked to the beta-APP allowing its expression to form beta-amyloid protein deposits in the animal's brain. The transgenic mouse is useful for elucidating the molecular mechanisms involved in the synthesis of and, more importantly, inhibiting the synthesis and deposition of beta-amyloid proteins (most importantly in the brain where plaque formation is associated with Alzheimer's disease) by inhibiting production and/or increasing cleavage after production. The transgenic animals provide useful models for studying the in vivo relationships of the proteins to each other and to other compounds being tested for their usefulness in treating Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-OCT-1994;
15-JUN-1990;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vaccines compositions for the treatment or prevention of infections by L. monocytogenes and related organisms.
Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APP; beta-amyloid precursor protein; human; transgenic mice; pathology; Alzheimer's disease; model; therapeutic compound; brain; mechanism; nerve tissue specific promoter; synthesis; inhibitor; deposition; plaque formation; treatment.
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Search completed: November 15, 2002, 11:44:12 Job time : 36 secs

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Copyright (c) 1993 - 2002 Compugen
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/cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/FCTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PcTUS_COMB.pep:*
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US-08-08-718-61
US-08-0443-976-61
US-08-443-977-61
US-09-499-302A-9
US-09-134-001C-5196
US-09-134-001C-5196
US-08-358-160-111
US-08-358-160-111
US-08-878-474-9
US-08-878-474-9
US-08-8791-1158-27
US-08-791-1158-27
US-08-791-1158-25
US-08-484-9938-4
US-08-484-9938-4
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(without alignments)
37.829 Million cell updates/sec
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Sequence 112, App Sequence 14, Appl Patent No. 5223482 Patent No. 5223482 Patent No. 5466783 Sequence 2, Appl Sequence 61, Appl Sequence 61, Appl Sequence 115, Appl Sequence 111, App Sequence 111, App Sequence 7, Appl Sequence 7, Appl Sequence 27, Appl Sequence 27
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| ication US/0835 N: Arthur C: Parthur C: RERNAN, Sonia K. ERNAN, Solia K. I RIGHERE I ONI ENGINEERE I ON DATA: US/08/3 16-DEC-194 N: 514 N: | 20222222222222222222222222222222222222 |
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| VED KUNITZ HUMAN NEUTROPHIL ELASTASE 300 | Sequence 4, Appli Sequence 12, Appli Sequence 6, Appli Sequence 4, Appli Sequence 39, Appli Sequence 1, Appli Sequence 1, Appli Sequence 3, Appli Sequence 4, Appli Sequence 17, Appli Sequence 17, Appli Sequence 17, Appli Sequence 17, Appli |

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RESULT 3
5187153-22
;'Patent No. 5187153
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; MOLECULE TYPE: protein
US-08-358-160-112
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US-08-422-333-14
                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30 .
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/422,333
EILING DATE: 13-APR-1995
CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: Shearer, Peter R.
REGISTRATION NUMBER: 21900-28048.00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 966-1550
TELEPHONE: (415) 966-1550
                                                                                                                                                         Query Match
Best Local
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Best Local Similarity 63.6%;
Matches 7; Conservative
                                                                                                                                         Matches
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                                                                                                                                                                                                                                                             TELEFAX: (415) 968-2438
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 60 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
TELEX: 24863
INFORMATION FOR SEQ ID NO:
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APPLICANT: CORDEL
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: TR
TITLE OF INVENTION: TH
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
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LENGTH: 60 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: sin
                                                                                                                                           Local Similarity es 7; Conserv
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F: 2450 Bayshore Parkway
Mountain View
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VENTION: TRANSGENIC NON-HUMAN MAMMAL DISPLAYING
VENTION: THE AMYLOID-FORMING PATHOLOGY OF ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
                                                                                                                                           Conservative
                                                                                                                                                                                                                     SS: single
linear
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linear
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Pred. No.
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Pred. No.
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RESULT 4
5220013 - 25
5220013 - 25
; Patent No. 5220013
; Patent No. 5220013
; PAPLICANT: PONTE, PHYLLIS A
; TITLE OF INVENTION: DNA SEQ:
; OF ALZHEIMER'S DISEASE
; NUMBER OF SEQUENCES: 30
; CURRENT APPLICATION DATA;
; CURRENT APPLICATION DATA;
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5220013-25
                                                                         RESULT 5
5223482-24
; Patent No. 5223482
; APPLICANT: SCHILLING,
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BARBARA
TITLE OF INVENTION: RECOMBINANT ALZHEIMER'S PROTEASE; INHIBITORY AMYLOID PROTEIN AND METHOD OF USE
NUMBER OF SEQUENCES: 34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5187153-22
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                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO:25:
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Best Local Similarity
Matches 7; Conserv
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PILING DATE: 30-NOV-989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 67,002
FILING DATE: 18-AUG-1987
APPLICATION NUMBER: 80,810
FILING DATE: 30-JAN-1987
APPLICATION NUMBER: 948,376
FILING DATE: 31-DEC-1986
APPLICATION NUMBER: 932,193
FILING DATE: 17-NOV-1986
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 361,912
FILING DATE: 06-JUN-1989
APPLICATION NUMBER: 359,911
FILING DATE: 12-MAY-1989
APPLICATION NUMBER: 87,002
FILING DATE: 18-AUG-1987
APPLICATION NUMBER: 8,810
FILING DATE: 18-AUG-1987
APPLICATION NUMBER: 8,810
FILING DATE: 30-JAN-1987
APPLICATION NUMBER: 948,376
FILING DATE: 31-DEC-1986
APPLICATION NUMBER: 932,193
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N: DNA SEQUENCE USEFUL FOR THE DETECTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        39.2%;
63.6%;
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63.6%;
                                                                            JAMES W.; PONTE,
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Pred. No.
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Pred. No.
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                                                                            PHYLLIS A.; CORDELL,
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35;
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RESULT 7
US-09-632-021-2
; Sequence 2, Application US/09632021
; Patent No. 6406896
; Patent No. 6406896
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Best Local Similarity
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TITLE OF INVENTION: HUMAN TISSUE FACTOR INHIBITOR

NUMBER OF SEQUENCES: 26

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO:5:
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                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local S
Matches 7
SEQ ID NO 2
                                                                      APPLICANT: Naumann, Todd A
TITLE OF INVENTION: Transposase Enzyme and Method for Use
FILE REPERIOE: 960296,96471
CURRENT APPLICATION NUMBER: US/09/632,021
CCURRENT FILING DATE: 2000-08-02
                             PRIOR APPLICATION NUMBER:
PRIOR FILING DATE: 1999-C
NUMBER OF SEQ ID NOS: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/07/36
FILING DATE: 06-JUN-1989
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 359,911
FILING DATE: 12-MAY-1989
APPLICATION NUMBER: 87,002
FILING DATE: 18-AUG-1987
APPLICATION NUMBER: 8,810
FILING DATE: 30-JAN-1987
FILING DATE: 30-JAN-1987
                                                                                                                                                 APPLICANT:
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FILING DATE: 15-JUL-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Wun, Tze-Chein,; Kretzmer, Kuniko K.; Broze,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
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FILING DATE: 31-DEC-1986
APPLICATION NUMBER: 932,193
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 566,280 FILING DATE: 13-AUG-1990
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                 Patentin Ver.
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                                              1999-08-02
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Best Local Similarity
Best Local Similarity
Best Local Similarity
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US-09-632-021-2
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                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: DK 4501/87
FILING DATE: 28 AUG-1987
PRIOR APPLICATION DATA:
APPLICATION UNMBER: DK 2254/88
FILING DATE: 26-APR-1988
PRIOR APPLICATION DATA:
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TYPE: PRT
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COMPUTER READABLE FORM:
REDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                               FILING DATE: 12-JUN-
PRIOR APPLICATION DATA:
                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: U
FILING DATE: 21-JUN-1
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE: 28-AUG-PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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                                                                                                                                     APPLICATION NUMBER: DK 2361/90 FILING DATE: 01-OCT-1990 PRIOR APPLICATION DATA:
                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                                                                                     APPLICATION NUMBER: US 0' FILING DATE: 19-NOV-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US FILING DATE: 19930623 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                 APPLICATION NUMBER: PCT/DK89/00096 FILING DATE: 25-APR-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
APPLICATION NUMBER: PCT/DK91/00299 FILING DATE: 01-OCT-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: No. 5591603o No. 5591603disk of No. 5591603th America, Inc. 405 Lexington Avenue, 62nd Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            United States of America
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No. 5591603rskov-Lauritsen, Leif
Christensen, Niels D.
Bregengaard, Claus
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No. 5591603ris, Kjeld
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                                                                                                                                                                                                                                                                                                                     NUMBER: PCT/DK88/00138
28-AUG-1988
                                                                                                                                                                                                               JMBER: US 07/466,408
21-JUN-1990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Process for the Preparation of Aprotinin and Aprotinin Analogs
                                                                                                                        US 07/598,337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US/08/084,718
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Pred. No. 2
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Best Local Similarity
Tatches 7; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-443-976-61
                                                                                                                                                       TELEPHONE: 212-867-0123
TELEFAX: 212-867-0298
INFORMATION FOR SEQ ID NO: 61:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 26-FEB-1993
ATTORNEY/AGENT INFORMATION:
NAME: Agris, Cheryl H.
REGISTRATION NUMBER: 34,086
REFERENCE/DOCKET NUMBER: 3143
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Christensen, Niels D.
APPLICANT: Bregengaard, Claus
APPLICANT: No. 561891513, Fanny
APPLICANT: Petersen, Lars C.
TITLE OF INVENTION: Process for the Preparation of Aprotinin
TITLE OF INVENTION: and Aprotinin Analogs
NUMBER OF SEQUENCES: 83
CORRESPONDENCE ADDRESS:
                                                                              APPLICATION NUMBER:
FILING DATE: 28-AUG-
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 111 amino acids
TYPE: amino acid
                                  PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US 07/827,687
FILING DATE: 29-JAN-1992
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                                                                                                                                                                                                                                                                                                                                        STATE: New York
COUNTRY: United States of America
ZIP: 10174-6201
                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: NO. 56189150 NO. 5618915disk of NO. 5618915th America, Inc. STREET: 405 Lexington Avenue, 62nd Floor CITY: New York
                                                APPLICATION NUMBER: DK 2254/88 FILING DATE: 26-APR-1988
                                                                                                                                APPLICATION NUMBER: US 08/084,718 FILING DATE: 23-JUN-1993
 APPLICATION NUMBER: FILING DATE: 28-AU
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o. 5618915
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Diness, Viggo
No. 5618915rskov-Lauritsen, Leif
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MBER: PCT/DK88/00138
28-AUG-1988
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26-FEB-1993
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                                                                                                                  DK 4501/87
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                US-08-443-977-61
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Best Local Similarity
Matches 7; Conserv
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                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                           tent No.
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                     APPLICANT: Peterser
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 111 amino acids
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FILING DATE: 26-FEB-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                     CORRESPONDENCE ADDRESS
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FILING DATE: 29-JAN-1
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 1
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APPLICATION NUMBER: DK 1118/91
FILING DATE: 12-JUN-1991
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PCT/DK89/00096
                                                                                                                                                       NUMBER OF SEQUENCES:
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APPLICATION NUMBER:
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APPLICATION NUMBER: DK 2361/90
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                                   COUNTRY: United States of ZIP: 10174-6201
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REGISTRATION NUMBER: 34,
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY:
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                                                                                                                   ADDRESSEE:
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                                                                                    New York
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                                                                  New York
                                                                                                405 Lexington Avenue,
                                                                                                                                                   Petersen, Lars C.

Petersen, Lars C.

VENTION: Process for the Preparation of Aprotinin

VENTION: and Aprotinin Analogs

EQUENCES: 83
                                                                                                                                                                                                                    Christensen, Niels D. Bregengaard, Claus No. 5621074ris, Fanny
                                                                                                                                                                                                                                                                          No. 5621074rskov-Lauritsen, Leif
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No. 5621074ris, Kjeld
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                                                                                                    No. 56210740 No. 5621074disk of No. 5 Lexington Avenue, 62nd Floor
Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UMBER: US 07/827,687
29-JAN-1992
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Pred. No. 84;
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                                                                                                                   5621074th America, Inc
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OPERATING SYSTEM:

IBM PC compatible

PC-DOS/MS-DOS

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RESULT 11
US-09-499-302A-9
                                                                                                                                                                                                                                                                     ; MOLECULE TYPE: US-08-443-977-61
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GENERAL INFORMATION:
APPLICANT: BOUNG-JUI
APPLICANT: MOON, K
                                             Sequence 9, Application US/09499302A Patent No. 6369212
                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: 212-867-012
TELEFAX: 212-867-0298
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 31
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 29-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER
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APPLICATION NUMBER: U
FILING DATE: 21-JUN-1
PRIOR APPLICATION DATA:
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FILING DATE: 28-AUG-1988
PRIOR APPLICATION DATA:
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FILING DATE: 23-JUN-1993
APPLICATION NUMBER: DK 4501/87
FILING DATE: 28-AUG-1987
PRIOR APPLICATION DATA:
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FILING DATE: 26-FEB-1993
ATTORNEY/AGENT INFORMATION:
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PRIOR APPLICATION DATA:
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APPLICATION NUMBER:
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FILING DATE: 25-APR-1989
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                                                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Agris, Cheryl H. REGISTRATION NUMBER: 3
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                                                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
                                                                                                                                                                           5 PCGAVMILYFH 15
                                                                                                                                            PCKARIILYFY 76
MOON, KYUNG KO
                                                                                                                                                                                                                                                                                                                                        111 amino acids
                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                        212-867-0123
                                                                                                                                                                                                                                                                                         protein
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                                                                                                                                                                                                                         38.2%;
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Pred. No.
                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                         DB 1;
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US-08-358-160-115
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.1 SEQ ID NO 9
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TITLE OF INVENTION: A CYTOCHROME P450 GENE HIGHLY EXPRESSED IN
TITLE OF INVENTION: INCOMPATIBLE INTERACTION
FILE REFERENCE: 10324/P64443USO
CURRENT FAPPLICATION NUMBER: US/09/499,302A
CURRENT FILLING DATE: 2000-02-07
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (243)
OTHER INFORMATION: (246)
NAME/KEY: MOD_RES
LOCATION: (246)
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OTHER INFORMATION: variable or unknown amino
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TYPE: PRT
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APPLICANT:
APPLICANT:
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NAME/KEY: MOD_RES
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OTHER INFORMATION:
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NAME/KEY: MOD_RES
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                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                              APPLICANT:
                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK
             SOFTWARE: PatentIn Release #1.0, Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                               TITLE OF INVENTION: ENGINEERED HUMAN-DERIVED KUNITZ
TITLE OF INVENTION: DOMAINS THAT INHIBIT HUMAN NEUTROPHIL ELASTASE
                                                                                                                                                                                                                                                                                 APPLICANT:
                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   459 GLLMLLYFFDW 469
                                                                                                                                                               CITY: Washington
                                                                                                                              COUNTRY:
                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity 54.
APPLICATION NUMBER:
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                                                                                                               20004
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                                                                                                                                                                             419 Seventh Street, N.W. Suite 300
                                                                                                                                                                                                                                                                                           GUTERMAN, Sonia K.
ROBERTS, Bruce L.
MARKLAND, William
                                                                                                                                                                                                                                                                                                                                                   LADNER,
                                                                                                                                USA
                                                                                                                                                                                                                                                                                 KENT, Rachel B.
                                                                                                                                                                                                                                                                                                                                                                  Arthur C.
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US/08/358,160
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Pred. No.
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PRIOR APPLICATION DATA:

16-DEC-1994

APPLICATION NUMBER:

FILING DATE:

13-OCT-1993

US 08/133,031

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                                                                                                                                    ; NUMBER OF SEQ ID NOS: 5674
; SEO ID NO 5196
; LENGTH: 346
; TYPE: PRT
; ORGANISM: Staphylococcus epidermidis
US-09-134-001C-5196
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1-09-134-001C-5196
                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Lynn Doucette-Stamm et al
APPLICANT: INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC-007
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 7; Conservat
                                                                                     Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 5196, Ap
Patent No. 638037
                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/055,779 PRIOR FILING DATE: 1997-08-14
                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 60/064,964 PRIOR FILING DATE: 1997-11-08
                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/134,001C CURRENT FILING DATE: 1998-08-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 202-737-3528
TELEX: 248633
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE CHARACTERISTICS:
LENGTH: 69 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/240,160
FILING DATE: 02-SEP-1988
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
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APPLICATION NUMBER: US 07/487,063
FILING DATE: 02-MAR-1990
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APPLICATION NUMBER:
129
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18 PCKAKMIRYFY 28
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REFERENCE/DOCKET NUMBER: LEY-1
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                                                                                   Local
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                             3 SZPCGAVMILYFHK-W 17
SIPLGIIAAVYYHKIW 144
                                                                   Similarity 7; Conservat
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                                                                   Conservative
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26-JAN-1993
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                                                                                   36.8%;
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                                                                   Score 37.5; DB 4;
Pred. No. 3.7e+02;
3; Mismatches 5;
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RESULT 14
US-08-358-160-110
; Sequence 110, A
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TELEX: 248633
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 58 amino acids
TYPE: amino acid
                                                              Best Local Similarity
Matches 7; Conserv
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                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/358,160
FILING DATE: 16-DEC-1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/133,031
FILING DATE: 13-OCT-1993
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US 07/664,989
FILING DATE: 01-MAR-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/487,063
FILING DATE: 02-MAR-1990
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                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/240,160
FILING DATE: 02-SEP-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 26-JAN-1993 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: KENT, Rachel B.
TITLE OF INVENTION: ENGINEERED HUMAN-DERIVED KUNITZ
TITLE OF INVENTION: DOMAINS THAT INHIBIT HUMAN NEUTROPHIL ELASTASE
NUMBER OF SEQUENCES: 234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
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                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
13
                                                                                                                                                                      TOPOLOGY: lin
                                                                                                                                                                                                                                                                            TELEPHONE: 202 - 3528
                                                                                                                                                                                                                                                                                                                                                           NAME: Cooper, Iver P. REGISTRATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Washington
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                               5 PCGAVMILYFH 15
PCKARMIRYFY 23
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419 Seventh Street, N.W. Suite 300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MARKLAND, William
                                                                 Conservative
                                                                                                                                                                        linear
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                                                                                 36.3%;
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                                                                 Score 37; DB Pred. No. 91; 1; Mismatches
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                                                                               DB 1; Length 58; 91;
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; TOPOLOGY: 1i
; MOLECULE TYPE:
US-08-358-160-111
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Search completed: November 15, 2002, 11:45:31 Job time : 15 secs
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-POS
SOSTWARE: PATENTION DATA:
APPLICATION STATE: 16-DEC-1994
CLASSIFICATION UMBER: US/08/358,160
FILING DATE: 16-DEC-1994
CLASSIFICATION UMBER: US 08/133,031
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/133,031
PILING DATE: 13-OCT-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/664,989
FILING DATE: 26-JAN-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/664,989
FILING DATE: US 07/664,989
FILING DATE: US 07/487,063
FILING DATE: US 07/487,063
FILING DATE: US 07/487,063
FILING DATE: US 07/487,063
FILING DATE: US-ARR-1990
PRIOR APPLICATION NUMBER: US 07/240,160
FILING DATE: US-SEP-1988
ATTORNEY/AGENT INFORMATION:
NAME: COOPET, IVET P.
REGISTRATION NUMBER: 28,005
REGISTRATION SUMBER: 28,005
REGISTRATION SUMBER: 28,005
REGISTRATION SUMBER: LEY-1
TELEPOMUNICATION SUMBER: 28,005
REGISTRATION SUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 36.3%;
Best Local Similarity 63.6%;
Matches 7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 59 amino acids
TYPE: amino acid
STRANDEDNESS: Single
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APPLICANT: GUTERMAN, Sonia K.
APPLICANT: ROBERTS, Bruce L.
APPLICANT: MARKLAND, William
APPLICANT: KENT, Rachel B.
TITLE OF INVENTION: ENGINEERED HUMAN-DERIVED KUNITZ
TITLE OF INVENTION: DOMAINS THAT INHIBIT HUMAN NEUTROPHIL ELASTASE
NUMBER OF SEQUENCES: 234
CORRESPONDENCE ADDRESS:
ANDRESCEP: TOTAL THE PROPERTY OF THE P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
ZIP: 20004
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STATE: District o
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419 Seventh Street, N.W. Suite 300
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length: 2000000000
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Gapop 10.0 , Gapext 0.5
  US-09-881-569A-2
102
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Published_Applications_AA:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BTSZPCGAVMILYFHKWR 18
                                                                                                                                                                                                                                                                                                                                                                                                       : /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*
: /cgn2_6/ptodata/2/pubpaa/US09_PUBCOMB.pep:*
: /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
: /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*
1: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
1: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
2: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
3: /cgn2_6/ptodata/2/pubpaa/US60_NEW_PUB.pep:*
4: /cgn2_6/ptodata/2/pubpaa/US60_NEW_PUB.pep:*
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/cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*
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/cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*
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10 US-09-815-242-11466

10 US-09-864-761-37111

12 US-10-028-051-8

9 US-09-903-170C-7

10 US-09-903-180B-7

10 US-09-903-187A-7

10 US-09-903-171A-7

10 US-09-903-171A-7

10 US-09-903-171A-7

10 US-09-903-1725A-7

10 US-09-903-323A-7

10 US-09-903-325A-7

9 US-09-903-170C-9

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9 US-09-903-180B-9
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10 US-09-881-569A-1

10 US-09-906-453-4

10 US-09-964-761-48038

10 US-09-764-870-484

10 US-09-895-913A-246
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27.110 Million cell updates/sec
Sequence 48038, A
Sequence 484, App
Sequence 21466, A
Sequence 37111, A
Sequence 7, Appli
Sequence 9, Appli
Sequence 9, Appli
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Sequence
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| 35 | ω 5 | 35 | 35 | 36 | 36 | 36 | 36 | 36 | 36 | 36 | 36 . | 36 | 36.5 | 36.5 | 36.5 | 36.5 | 36.5 | 37 | 37 | 37 | 37 | 37 | 37 | 37 | 37 |
| 34.3 | 34.3 | 34.3 | 34.3 | 35.3 | 35.3 | 35.3 | 35.3 | 35.3 | 35.3 | 35.3 | 35.3 | 35.3 | 35.8 | 35.8 | 35.8 | 35.8 | 35.8 | 36.3 | 36.3 | 36.3 | 36.3 | 36.3 | 36.3 | 36.3 | 36.3 |
| 160 | 136 | 66 | 38 | 1356 | 891 | 842 | 639 | 552 | 276 | 57 | 49 | 28 | 267 | 262 | 262 | 262 | 71 | 403 | 325 | 325 | 325 | 325 | 325 | 325 | 325 |
| 10 | 10 | 10 | 10 | 10 | 10 | 10 | 10 | 9 | 10 | œ | 10 | 10 | 10 | 10 | 10 | 10 | 12 | 10 | 12 | 12 | 10 | 10 | 10 | 10 | 10 |
| US-09-731-872-316 | US-09-970-711-5 | US-09-764-877-1085 | US-09-864-761-44559 | US-09-801-368-306 | US-09-949-192-45 | US-09-798-831-8 | US-09-768-877-25 | US-09-843-676-4 | US-09-925-300-1624 | US-08-896-322-5 | US-09-764-887-204 | US-09-864-761-44551 | -09-925 | US-09-988-975A-7 | US-09-764-762-4 | -09 | US-10-001-879-194 | US-09-870-379-2 | US-10-028-051-4 | US-10-028-051-2 | US-09-903-325A-9 | -903-323A | US-09-903-188A-9 | -09-903- | US-09-903-187A-9 |
| Sequence 316, App | Sequence 5, Appli | Sequence 1085, Ap | Sequence 44559, A | 5 | Sequence 45, Appl | Sequence 8, Appli | Sequence 25, Appl | Sequence 4, Appli | Sequence 1624, Ap | Sequence 5, Appli | Sequence 204, App | Sequence 44551, A | Seguence 745, App | Sequence 7, Appl1 | Sequence 4, Appl1 | Sequence 9, Appli | Sequence 194, App | Sequence 2, Appli | • | Ņ | • | • | • | Sequence 9, Appli | • |

ALIGNMENTS

RESULT 1 US-09-881-569A-2

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Sequence 2, Application US/09881569A

Patent NO. US20020151867A1

GENERAL INFORMATION:
APPLICANT: KHANNA, PUSHPA

ITLE OF INVENTION: AND A PROCESS FOR THE EXTRACTION THEREOF

FILE REFERENCE: 013488-3

CURRENT APPLICATION UNMBER: US/09/881,569A

CURRENT APPLICATION UNMBER: 560/DEL/99

PRIOR APPLICATION NUMBER: 560/DEL/99

PRIOR FILING DATE: 1999-04-13

ESULT 2
US-09-801-569A-1
Sequence 1, Application US/09881569A
Patent No. US20020151687A1
Patent No. US20020151687A1
GENERAL INFORMATION:
APPLICANT: KHANNA, PUSHPA
TITLE OF INVENTION: PROTEIN/POLYPEP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Description of Artificial Sequence: ; OTHER INFORMATION: isolated from momordica charantia US-09-881-569a-2
                                                                                                                                                                                                                                         밁
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LENGTH: 18

TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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Best Local (
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                                                                                                                                                                                                                                                                                                                                              Local Similarity 
nes 18; Conserv
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     PROTEIN/POLYPEPTIDE-K OBTAINED FROM MOMORDICA CHARANTIA
                                                                                                                                                                                                                                                                                                                                                                      100.0%;
                                                                                                                                                                                                                                                                                                                                                 0
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Pred. No. 3.4e-10;
; Mismatches 0;
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CURRENT APPLICATION NUMBER: US/09/881,569A
CURRENT FILING DATE: 2001-06-14
PRIOR APPLICATION NUMBER: 560/DEL/99
PRIOR FILING DATE: 1999-04-13
PRIOR APPLICATION NUMBER: 561/DEL/99
PRIOR FILING DATE: 1999-04-13
PRIOR FILING DATE: 1999-04-13
NUMBER OF SEQ ID NOS: 2
SEQ ID NO 1
SEQ ID NO 1
LENGTH: 16
                                                                                                                                                                                             US-09-864-761-48038; Sequence 48038, Application US/09864761; Patent No. US20020048763A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Zea mays
US-09-906-453-4
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                                                                                                                               GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: PatentIn Ver. SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4, Application US/09906453 Patent No. US20020120125A1
                                                 APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
FILE REFERENCE: Acomica-x-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Polycomb genes from Malze - FILE REFERENCE: WISCONSINIOOUS CURRENT APPLICATION NUMBER: US/09/906,453 CURRENT FILING DATE: 2001-07-16
                CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Helentjaris, Timothy APPLICANT: Phillips, Ronald
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Kaeppler, Shawn APPLICANT: Springer, Nath
PRIOR APPLICATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQ ID NOS: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: AND FILE REFERENCE: 013488-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                   317 ENKKPCGHLCYLRFPQWR 334
                                                                                                                                                                                                                                                                                                                                  1 BTSZPCGAVMILYFHKWR 18
                                                                                                                                                                                                                                                                                                                                                                      Local Similarity es 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                      Conservative
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 NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Description of Artificial Sequence: Polypeptide-p isolated from momordica charantia
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38.98;
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62.5%;
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Pred. No.
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18;
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0.0018;
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                                                                                                                                                                                                                                                                                                                                                                                                   Length 893;
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; ORGANISM: Homo sapiens US-09-764-870-484
                                                                                                                                                                                                                                                                                                        RESULT 5
US-09-764-870-484
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                                     Prior application data removed -
NUMBER OF SEQ ID NOS: 646
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 484
LENGTH: 74
TYPE: PRT
                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins,
FILE REFERENCE: PT214
                                                                                                                                                                                                                                                                     Sequence 484, Application US/09764870 Patent No. US20020042386A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: Annomax Sequence Listing Engine vers. 1.1 SEQ ID NO 48038
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/764,870 CURRENT FILING DATE: 2001-01-17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                           3 SZPCGAVMILYFHKW 17
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Pred. No.
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; APPLICANT: Xu, H. Howard
; TITLE OF INVENTION: Identification of Essential Genes in ITTLE OF INVENTION: Prokaryotes
; TITLE OF INVENTION: Prokaryotes
; FILE REFERENCE: ELITRA.011A
; CURRENT APPLICATION NUMBER: US/09/815,242
; CURRENT APPLICATION NUMBER: 60/191,078
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR APPLICATION NUMBER: 60/206,848
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US-09-815-242-11466
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SEQ ID NO 246
LENGTH: 1185
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APPLICANT: Oomen, Raymond P.
TITLE OF INVENTION: Identification of Polynucleotides
TITLE OF INVENTION: Encoding No. US20020160456A1el Helicobacter Polypeptides in the
TITLE OF INVENTION: Genome
TILE REFERENCE: 06132/043002
CURRENT APPLICATION NUMBER: US/09/895,913A
CURRENT FILING DATE: 2001-06-29
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APPLICANT:
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ORGANISM: Helicobacter pylori
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5 TSRKCGAVVII 15
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                                  APPLICATION NUMBER: 60/242,578
 APPLICATION NUMBER:
                  FILING DATE:
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o. US20020160456A1
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Trawick, John D.
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Zyskind, Judith W.
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Tomb, Jean François
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60/253,625
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Pred. No.
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US-09-864-761-37111
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PRIOR FILING DATE: 2000-12-2
PRIOR APPLICATION NUMBER: 60/269,308
PRIOR FILING DATE: 2001-02-16
NUMBER OF SEQ ID NOS: 14110
SOFTWARE: FastSEQ for Windows Version
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CURRENT APPLICATION NUMBER: US/09/864,761
CURRENT FILING DATE: 2001-05-23
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APPLICANT: Chen, Wensheng

TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC

TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
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TYPE: PRT
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PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR FILING DATE: 2000-09-21
PRIOR APPLICATION NUMBER: US 09/608,408
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PRIOR APPLICATION NUMBER: PCT/US01/00667
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PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/00666
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PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR FILING DATE: 2000-08-03
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                                            PRIOR FILING DATE:
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RAPPLICATION NUMBER: PCT/USO1/00663
RAPPLICATION DATE: 2001-01-30
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APPLICATION NUMBER: PCT/US01/00669
FILING DATE: 2001-01-30
                                          FILING DATE: 2000-06-30
APPLICATION NUMBER: US
FILING DATE: 2001-01-29
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                        ID NOS:
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RESULT 9
US-10-028-051-8
; Sequence 8, Application US/10028051
; Patent No. US20020147329A1
; GENERAL INFORMATION:
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Best Local S
Matches 6
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OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 1.2
OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 9.2
OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 10
OTHER INFORMATION: EXPRESSED IN HEAL SIGNAL = 9.2
OTHER INFORMATION: EXPRESSED IN HEAL SIGNAL = 12
OTHER INFORMATION: EXPRESSED IN HELA SIGNAL = 12
OTHER INFORMATION: EXPRESSED IN HELA SIGNAL = 10
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 10
OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 13
OTHER INFORMATION: EXPRESSED IN HIT: AW411290.1, EVALUE 5.00e-05
OTHER INFORMATION: SWISSPROT HIT: P81274, EVALUE 9.00e-06
                                                                                                                         TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: MAP TO OTHER INFORMATION: EXPRESS OTHER INFORMATION: EXPRESS
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les 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/028,051
FILING DATE: 19-Dec-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: BATLfeld, Neil S
REGISTRATION NUMBER: 39,901
REFERENCE/DOCKET NUMBER: NIH133.001A
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-235-8550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Knobbe, Martens, Olson & Bear
STREET: 620 Newport Center Drive, 16th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Luyten,
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COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ Version 1.5
TYPE: amino a STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 08/729,452 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 620 Newport CITY: Newport Beach
                                     TYPE: amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A.
ZIP: 92660
                                                                                                                                                                                                                                                                      TELEFAX: 619-235-0176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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42.9%;
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Pred. No.
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31;
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APPLICANT: De Robertis, Edward M.
APPLICANT: Bouwmeester, Tewis
TITLE OF INVENTION: Endoderm, Cardiac and Neural Inducing
TITLE OF INVENTION: Factors
FILE REFERENCE: 510015-259
CURRENT APPLICATION NUMBER: US/09/903,170C
CURRENT APPLICATION NUMBER: US 60/020,150
PRIOR APPLICATION NUMBER: US 60/020,150
PRIOR FILING DATE: 1996-06-20
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 7
LENGTH: 323
TYPE: PRT
ORGANISM: Mouse FRZB-1
US-09-903-170C-7
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                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: De Robertis, Edward M.
APPLICANT: Bouwmeester, Tewis
TITLE OF INVENTION: Endoderm, Cardiac and Neural Inducing
TITLE OF INVENTION: Factors
FILE REFERENCE: 510015-256
CURRENT APPLICATION NUMBER: US/09/903,180B
CURRENT FILING DATE: 2001-07-11
                                                                                                                                                   SEQ ID NO 7
LENGTH: 323
TYPE: PRT
ORGANISM: Mouse FRZB-1
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 7, Application US/09903180B Patent No. US20020099171A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 35.
Matches 5; Conservative
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Best Local Similarity 35.7%;
Matches 5; Conservative
                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: US 60/020,150 PRIOR FILING DATE: 1996-06-20 NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                         SOFTWARE: FastSEQ for Windows Version 3.0
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117 QGCEPILIKYRHSW
                                   4 ZPCGAVMILYFHKW 17
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                                                                    Conservative
                                                                                    36.38;
35.78;
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                                                                                    Score 37;
Pred. No.
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Pred. No.
                                                                    Mismatches
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                                                                                    DB
85;
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84;
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                                                                                                  Length 323;
                                                                    Indels
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RESULT 12 US-09-903-187A-7

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LENGTH: 323
TYPE: PRT
ORGANISM: Mouse
US-09-903-187A-7
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SEOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 7
LENGTH: 323
TYPE: PRT
ORGANISM: Mouse FRZB-1
US-09-903-171A-7
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Best Local Similarity
Watches 5; Conserve
                                                RESULT 14
US-09-903-188A-7
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GENERAL INFORMATION:
             Sequence 7, Application US/09903188A Patent No. US20020128439A1
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Best Local :
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 7
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APPLICANT: Bouwmeester, Tewis
TITLE OF INVENTION: Endoderm, Cardiac and Neural Inducing
TITLE OF INVENTION: Factors
FILE REFERENCE: 510015-248
                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/903,171A CURRENT FILING DATE: 2001-07-11 PRIOR APPLICATION NUMBER: US 60/020,150 PRIOR FILING DATE: 1996.06-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: De Robertis, Edward M.
APPLICANT: Bouwmeester, Tewis
TITLE OF INVENTION: Endoderm, Cardiac and Neural Inducing
TITLE OF INVENTION: Factors
FILE REFERENCE: 510015-260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: 09/552,988
PRIOR FILING DATE: 2000-04-21
PRIOR APPLICATION NUMBER: US 08/878,474
PRIOR FILING DATE: 1997-06-18
PRIOR APPLICATION NUMBER: US 60/020,150
PRIOR FILING DATE: 1996-06-20
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                                                                                                                      117 QGCEPILIKYRHSW 130
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35.7%;
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Pred. No. 85;
3; Mismatches
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US-09-903-323A-7
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US-09-903-188A-7
Job time
               Search completed: November 15,
                                                                      В
                                                                                                                                                                                                                      US-09-903-323A-7
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LENGTH: 323
TYPE: PRT
                                                                                                                                                                                                                                    SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 7
LENGTH: 323
TYPE: PRT
ORGANISM: Mouse FRZB-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 7, Application US/09903323A Patent No. US20020128440A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 35.7%;
Matches 5; Conservative
                                                                                                                                              Matches
                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Endoderm, Cardiac and Neural Inducing FILTE OF INVENTION: Factors FILE REPERENCE: 510015-561
CURRENT APPLICATION NUMBER: US/09/903,323A
CURRENT FILING DATE: 2001-07-11
PRIOR APPLICATION NUMBER: US 60/020,150
PRIOR APPLICATION NUMBER: US 60/020,150
PRIOR FILING DATE: 1996-06-20
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: De Robertis, Edward M. APPLICANT: Bouwmeester, Tewis
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APPLICANT: Bouwmeester, Tewis
TITLE OF INVENTION: Endoderm, Cardiac and Neural Inducing
TITLE OF INVENTION: Factors
FILE REFERENCE: 510015-258
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                                                                      117 QGCEPILIKYRHSW
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| multiple sugar-bin | F70018 | N | 422 | 38.2 | 39 | σ |
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| flavodoxin homolog | D69541 | N | 148 | 38.2 | 39 | N |
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| ATP-dependent deox | AD1358 | N | 1157 | 38.7 | 39.5 | 0 |

ALIGNMENTS

Fleischmann, R.D.; Quackenbush, J.F.; White, O.; Nelson, K.E.; Ketchum, K.A.; Down, Fleischmann, R.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, 1 Glodek, A.; Zhou, L.; Overbeek, R.; Gocayne, J.D.; Weidman, J.F.; McDonald, L. Nature 390, 364-370, 1997

A;Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Kaine, B.P.; Sykes, A;Authors: The complete genome sequence of the hyperthermophilic, sulfate-reducing arch A;Reference number: A69250; MUID:98049343; PMID:9389475

A;Accession: B69433

A;Status: Traisgamma-glutamyl carboxylase (EC 4.1.1.-) - bovine C;Species: Bos prinigenius taurus (cattle) C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 05-Nov-1999 C;Date: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 05-Nov-1999 C;Accession: A47439; A38939; B39283; C39283 R;Rehemtulla, A.; Roth, D.A.; Wasley, L.C.; Kuliopulos, A.; Walsh, C.T.; Furie, Proc. Natl. Acad. Sci. U.S.A. 90, 4611-4615, 1993 A;Title: In vitro and in vivo functional characterization of bovine vitamin K-de A;Reference number: A47439; MUID:93281629; PMID:8506307 A;Accession: A47439 signal-transducing histidine kinase homolog - Archaeoglobus fulgidus (;Species: Archaeoglobus fulgidus C;Species: Archaeoglobus fulgidus C;Date: 05-Dec-1997 *sequence_revision 05-Dec-1997 *text_change 21-Jul-2000 C;Accession: .B69433 R;Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O; Nelson, K.E.; Ketchum, R;Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O; Nelson, K.E.; Ketchum, A;Cross-references: GB:L09726; NID:g289398; PIDN:AAA30425.1; PID:g289399 R;Wu, S.M.; Cheung, W.F.; Frazier, D.; Stafford, D.W. submitted to GenBank, January 1992 A;Status: preliminary A;Molecule type: mRNA A;Residues: 1-758 <REH> A;Status: preliminary; nucleic acid sequence not shown; translation A;Molecule type: DNA A;Residues: 1-222 <KLE> A;Cross-references: GB:M81593; NID:g162744; PIDN:AAA30410.1; PID:g162745 R;Wu, S.M.; Cheung, W.F.; Frazier, D.; Stafford, D.W. A; Molecule type: mRNA A; Residues: 96-758 <WUA> A; Reference number: A38939 A; Accession: A38939 A;Cross-references: GB:AE001001; GB:AE000782; NID:g2689324; PIDN:AAB89779.1; PID:g264 Query Match 204 NEPCGAVFVMEF 215 Local Similarity hes 6; Conserv 3 SZPCGAVMILYF 14 Conservative 43.18; 4; Mismatches Score 44; DB: Pred. No. 7.4; DB 2; 2; Length 222; Indels bovine vitamin K-depende 0 not shown Gaps 0

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A;Cross-references: GB:AE001259; GB:AE000520; NID:g3323209; PIDN:AAC65853.1; PID:g33232:A;Experimental source: strain Nichols C;Genetics;
                                                                                                                              rson, J.; Khalak, H.; Richardson, D.; Howell, J.K.; Chidambaram, M.; Uthey, L.; Weldman, J.; Smith, H.O.; Venter, J.C.
Science 281, 375-388, 1998
A;Title: Complete genome sequence of Treponema pallidum, the syphilis A;Reference number: A71250; MUID:98332770; PMID:965876
A;Accession: G71266
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nakazaki, N.; Shimpo, S.; Sugimoto, M.; Ta
DNA Res. 8, 205-213, 2001
A;Title: Complete Genomic Sequence of the I
A;Reference number: AB1807; MUID:21595285;
A;Accession: AC2277
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A;Title: Cloning and expression of the cDNA for human gamma-glutamyl carboxylase.
A;Reference number: A39283; MUID:92086858; PMID:1749935
A;Accession: B39283
A;Status: nucleic acid sequence not shown
                                                                                                                                                                                                                                                                      probable ATP-dependent nuclease, subunit A - syphilis spirochete
C;SpecLes: Treponema pallidum subsp. pallidum (syphilis spirochete)
C;Date: 24-Jul-1998 #sequence_revision 24-Jul-1998 #text_change 05-Nov-1999
C;Accession: G71266
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C; Keywords: carbon-carbon
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A; Residues: 37-50 <WU3>
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A;Accession: C39283
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A; Residues: 96-693,695-758
                                                                        A; Molecule type: DNA
A; Residues: 1-1239 <COL>
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R;Fraser, C.M.; Norris, S.J.; Weinstock, G.M.; White, O.;
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A; Residues: 1-64 < KUR>
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hypothetical protein asr3770 [imported] - Nostoc sp. (strain PCC 7120)
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set domain protein - Chlamydophila pneumoniae (strain CWL029) C;SpecLes: Chlamydophila pneumoniae, Chlamydia pneumoniae C;Date: 23-Apr-1999 #sequence_revision 23-Apr-1999 #text_chan: C;Accession: D72024
                                                                                       RESULT
D72024
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, C.; Dodson, R.; Gwinn, M.; Nelson, W.; DeBoy, R.;
Nucleic Acids Res. 28, 1397-1406, 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        conserved hypothetical protein CP0991 [imported] - Chlamydophila pneumoniae C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae C;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 11-May-2000 C;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 11-May-2000
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A;Title: Comparison of whole genome sequences of chlamydia A;Reference number: A86491; MUID:20330349; PMID:10871362 A;Accession: D86600
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C;Species: Chlamydophila pneumoniae, Chlamydia pneumoniae
C;Date: 02-Mar-2001 #sequence_revision 02-Mar-2001 #text_change 02-Mar-2001
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A; Residues; 1-221 <REA>
A; Cross-references: GB:AE002257; GB:AE002161;
A; Experimental source: strain AR39, HL cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Title: Genome sequences of Chlamydia trachomatis MoPn and Chlamydia pneumoniae A;Reference number: A81500; MUID:20150255; PMID:10684935
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A; Residues: 1-221 <STO>
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Pred. No. 52;
1; Mismatches
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Pred. No. 16;
5; Mismatches
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Salzbe
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C; Accession: H83589

R; Stover, C.K.; Pham, X.Q.; Erwin, A.L.; Mizoguchi, S.D.; Warrener, P.; Hickey, adman, S.; Yuan, Y.; Brody, L.L.; Coulter, S.N.; Folger, K.R.; Kas, A.; Larbig, acman, S.; Olson, M.V.
Nature 406, 959-964, 2000
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Best Local Similarity
""" 6; Conserv
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                                                                                                                                                                 A;Title: Complete genome sequence of Pseudomonas aeruginosa A;Reference number: A82950; MUID:20437337; PMID:10984043 A;Accession: H83589
                                                                                                                                                                                                                                                                                                                conserved hypothetical protein PA0451 [imported] - Pseudomonas aeruginosa (strain C;Speciles: Pseudomonas aeruginosa C;Date: 15-Sep-2000 #sequence_revision 15-Sep-2000 #text_change 31-Dec-2000 C;Accession: H83589
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A;Accession: AE0958
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A; Residues: 1-221 <ARN>
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A; Title: Comparative genomes of Clamydia pneumoniae and C. trachomatis.
A; Reference number: A72000; MUID:99206606; PMID:10192388
                                                            A;Cross-references: GB:AE004482; GB:AE004091; A;Experimental source: strain PAO1
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A; Residues: 1-443 <STO>
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;Superfamily: Rhizobium nodulation competitiveness
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                                           Genetics:
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Pred. No.
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Davis, P.; Davies, R.M.; Dowd, L.; White, N.;
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                                         C; Genetics:
A;Genetic
                                                            A; Experimental source: strain G-37
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A;Cross-references: EMBL:U39725; NID:g3845017; PID:g3845030
A;Experimental source: isolate G37
R;Fraser, C.M.; Gocayne, J.D.; White, O.; Adams, M.D.; Clayton, R.A.; Fleischmann, R. M.; Fuhrmann, J.; Nguyen, D.; Utterback, T.R.; Saudek, D.M.; Phillips, C.A.; Merrick, C.A.; Venter, J.C.
Science 270, 397-403, 1995
A;Title: The minimal gene complement of Mycoplasma genitalium.
A;Reference number: A64200; MUID:98026346; PMID:7589993
A;Accession: C64248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      phosphatidate cytidylyltransferase (EC 2.7.7.41) - Mycoplasma genitalium N;Alternate names: CDP-diglyceride synthetase C;Specles: Mycoplasma genitalium C;Specles: Mycoplasma genitalium C;Date: 16-Jul-1999 *sequence_revision 16-Jul-1999 *text_change 02-Jun-2000
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C;Species: Arabidopsis thaliana (mouse-ear cress)
C;Date: 02-Feb-2001 #sequence_revision 02-Feb-2001 #text_change 02-Feb-2001
                                                                                                                                                                                                                                                                                                                                                                                                submitted to the EMBL Data A; Reference number: Z16818 A; Accession: T09771
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R;Fraser, C.M.; Gocayne, J.D.; White, O.; Adams, M.D.; Clayton, R.A.; Fleischmann, R., M.; Fuhrmann, J.L.; Nguyen, D.T.; Utterback, T.; Saudek, D.M.; Phillips, C.A.; Merr
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A; Residues: 1-353 <STO>
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A;Cross-references: GB:U39730; GB:L43967; NID:g1046149;
                         A; Molecule type: DNA
A; Residues: 70-374 < TIGR>
                                                                          A; Status: nucleic acid sequence
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A; Residues: 1-374 <FRA>
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5; Conserv
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8; Conserv
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27.8%;
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                                                                          not shown;
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Pred. No.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    October 1998
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                                                                          translation not shown
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BD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
     PIDN: AAB01627.1; PID: g1046154
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A: SCBRUB: F.C...
A: Molecule type: DNA
A: Residues: 1-544 < PAN>
A: Cross • references: EMBL: Z95496;
A: Cross • references: CV. Igri
RESULT
AB1793
                                                                                                                                                                                             A;Description: catalyzes the hydrolysis of chitin, A;Pathway: chitin degradation C;Superfamily: Serratia marcescens chitinase C;Keywords: glycosidase; hydrolase; polysaccharide
                                                                                                                                                                                                                                                                              A;Molecule type: DNA
A;Residues: 1-558 <KUZ>
A;Cross-references: EMBL:AF081810; NID:g3822234; PIDN:AAC70256.1; PID:g3822305
                                                                                                                                                                                                                                                                                                                                                R;Kuzio, J.; Pearson, M.N.; Harwood, S.H.; Funk, C.J.; Evans, J.T.; Slavicek, Virology 253, 17-34, 1999
A;Title: Sequence and analysis of the genome of a baculovirus pathogenic for A;Reference number: Z20836; MUID:99124785; PMID:9887315
A;Accession: T30418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT
T30418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     submitted to the EMBL Data Library, May 1997 A;Reference number: Z15093 A;Accession: T05952
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Species: Hordeum vulgare (barley)
C;Date: 30-Apr-1999 #sequence_revision 30-Apr-1999 #text_change 15-Jun-2001
Accession: T05952
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A; Introns: 41/3; 1:
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                                                                   밁
                                                                                                                                                                                                                                                                                                                               A; Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                               chitinase (EC 3.2.1.14) - Lymantria dispar nuclear polyhedrosis virus

Species: Lymantria dispar nuclear polyhedrosis virus, LdMNPV

Date: 29-Oct-1999 #sequence_revision 29-Oct-1999 #text_change 21-Jul-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Gene: Mlo-hl
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Best Local
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                                                                                                                                               Local Similarity
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                                                                                               ω
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                                                                                             SZPCGAVMILYFHKW 17
                                                                   AEPTGKVVAAYFVEW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    10; Conservative
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                                                                                                                                  Conservative
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Pred. No.
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Pred. No. 38;
                                                                                                                                                               Score 41;
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Science 294, 849-852, 2001
A; Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J. A; Title: Comparative genomics of Listeria species.
A; Reference number: AB1077; MUID:21537279; PMID:11679669
A; Accession: AH1417
A; Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                               C;Date: 27-Nov-2001 #sequence_revision 27-Nov-2001 #tes
C;Accession: AH1417
R;Glaser, P.; Frangeul, L.; Buchrieser, C.; Amend, A.;
; Dominguez-Bernal, G.; Duchaud, E.; Durand, L.; Dussu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J. A;Title: Comparative genomics of Listeria species, A;Reference number: AB1077; MUID:21537279; PMID:11679669
A;Accession: AB1793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Date: 27-Nov-2001 #sequence_revision 27-Nov-2001 #tey
C;Accession: AB1793
R;Glaser, P; Frangeul, L.; Buchrieser, C.; Amend, A.;
Dominguez-Bernal, G.; Duchaud, E.; Durand, L.; Dussu
В
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                                                                                                                                      C; Superfamily:
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A; Residues: 1-575 <GLA>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 1-575 <GLA>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              D.; Jones, L.M.; Karst, U. Science 294, 849-852, 2001
                                                                                                                                                                                         A; Experimental source: strain
                                                                                                                                                                                                           A;Cross-references: GB:NC_003210; PIDN:CAD00958.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; Species:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABC transporter (ATP-binding protein) homolog lmo2745 [imported] - Listeria
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                                                                   Query Match
Best Local S
Matches 6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transporter (ATP-binding protein) homolog lin2888 [imported] - Listeria
150
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nes 6; Conserv
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GAILLEFFMDWK 161
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Pred. No.
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Mismatches
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surget, O.;
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surget, O.;
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1 BTSZPCGAVMILYFHKWR 18
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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| 0; 0; | is SWISS-PROT entry is copyrig tween the Swiss Institute of European Bioinformatics Inste by non-profit institution diffied and this statement is not titles requires a license agressend an email to license@isb- | Velazquez-Estades rat vitamin K-depe lly regulated gene lly regulated gene s. 243:334-346(199 GLUTAMIC ACID RES E RESIDUES TO GAMM MAY FORM A HOMODITY: SOME, TO PLANT | Dawley: TISSUE- B: PubMed-97040 B: PubMed-97040 COR., Velazquez tural organizat tural organizat K-dependent ga S. Res. Commun. | (Rat). ; Chordata; ; Rodentia; | 37, Created) 37, Last sec 37, Last anr 1t gamma-carl | TANDARD; | | 4429 4429 502 502 661 838 838 838 839 |
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"In vitro and in vivo functional characterization of I K-dependent gamma-carboxylase expressed in Chinese hau
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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ifled and this statement is not removed. Usage by an
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PTM: THE N-TERMINUS IS BLOCKED.
SIMILARITY: SOME, TO PLANT AND MAMMALIAN LIPOXYGENASES.
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STRAIN-ATCC 33530 / G-37;
STRAIN-94075230; PubMed-8253680;
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Therson S.N., Hu P.-C., Bott K.F., Hut-
Mycoplasma genitalium
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                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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-i- PATHWAY: Phospholipid biosynthesis.
-i- SUBCELLULAR LOCATION: Integral members
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    -1- SUBCELLULAR LOCATION: Integral membran
    -1- SIMILARITY: BELONGS TO THE CDS FAMILY.

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CATALYTIC ACTIVITY: CTP + phosphatidate
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U02189; AAD12473.1;
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                                                                                                         "A new Kunitz-type inhibitor from bovine determination.";
                                                                                                                                                                                                                  21-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Serum basic protease inhibitor (Serum BPI).
Bos taurus (Bovine).
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                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea Bovidae; Bos. Bos. NCBI_TaxID-9913;
        InterPro; IPR002223; Kunitz_BPTI;
Pfam; PF00014; Kunitz_BPTI; 1.
PRINTS; PR00759; BASICPTASE.
                                                                                                     FEBS Lett. 119:58-62(1980)
                                                                                                                                                         SEQUENCE
                                                                                                                                                                                                                                                                        IBPS_BOVIN P00975;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR004326; Mlo. Pfam; PF03094; Mlo; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (MAY-1997) to the EMBL/GenBank/DDBJ databases i- SUBCELLULAR LOCATION: Integral membrane protein (Po-1-SIMILARITY: BELONGS TO THE MLO FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Submitted (MAY-1997)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN-cv. Igri;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hordeum vulgare (Barley).
Eukaryota; Viridiplantae;
                                       PIR; A01206; TIBOR.
HSSP; P00974; 1BPI.
                                                           -!- SUBCELLULAR LOCATION: Secreted.
-!- SIMILARITY: CONTAINS 1 BPTI/KUNITZ
                                                                               -i- FUNCTION: THIS INHIBITOR HAS ACTIVITY VERY SIMILAR TO BASIC PROTEASE INHIBITOR FROM BOVINE TISSUES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; 295496; CAB08860.1;
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annotation update)
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YBO2_YE
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01-OCT-1994 (Rel. 30, I
15-DEC-1998 (Rel. 37, I
Hypothetical 11.5 kDa F
YBR032W OR YBR0317.
P47655;
01-FEB-1996
01-FEB-1996
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ACT_SITE
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P38223;
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DISULFID 7 57
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ACT_SITE 17 18
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                                                                                                                                                                                                                                                 Hypothetical SEQUENCE 10
                                                                                                                                                                                                                                                                           PIR; S45888; S45888.
SGD; S0000236; YBR032W.
                                                                                                                                                                                                                                                                                                                                      or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                 the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                              This
                                                                                                                                                                                                                                                                                                                                                                                                                                                      chromosome II from Saccharomyces cerevisiae reveals frames, including ten new open reading frames, five identified genes and a homologue of the SCOl gene. "; yeast 10:S75-S80(1994).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=94378725; PubMed=8091864;
Smits P.H.M., de Haan M., Maat C., Grivell L.A.;
"The complete sequence of a 33 kb fragment on the right arm of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota; Fungi; i
Saccharomycetales;
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33,
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01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
UDP-glucuronosyltransferase 1-3 precursor, microsom
UDP-glucuronosyltransferase (1-3) (UGT1A3) (B3).
                       MEDLINE=90274676; PubMed=2112380; Sato H., Kolwai O., Tanabe K., Ka "Isolation and sequencing of rat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a between the Swiss Institute of Bioinformatics and the EMBLE the European Bioinformatics Institute. There are no restrict use by non-profit institutions as long as its content is modified and this statement is not removed. Usage by and the entitles requires a license agreement (See http://www.isb-sib or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fritchman J.L., Weidman J.F., Small K.V., Sandusky M., Fuhrmann Nguyen D.T., Utterback T.R., Saudek D.M., Phillips C.A., Merrick Tomb J.F., Dougherty B.A., Bott K.F., Hu P.-C., Lucier T.S., Peterson S.N., Smith H.O., Hutchison C.A. III, Venter J.C.; "The minimal gene complement of Mycoplasma genitalium."; Science 270:397-403(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
STRAIN-ATCC 33530 / G-37;
MEDLINE-96026346; PubMed-7569993;
             glucuronosyltransferase
                                                                                                                                                            MEDLINE=95332265; PubMed=7608130; Emi Y., Ikushiro S.I., Iyanagi T.
                                                                                                                                                                                           STRAIN-Wistar;
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Mammalia; Eutheria;
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Bacteria; E
primary
                                                                               SEQUENCE OF
                                                                                                                                 Emi Y., Ikushiro S.I.
"Drug-responsive and
multiple first exons
                                                                                                                                                                                                      SEQUENCE OF 1-286 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                   064637;
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5; Conserv
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transcript.
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272 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 protein;
                                                                                                                                                                                                                                                                                                                                                                                STANDARD;
                                                                                                                                                                                                                                                                       (Rat).
                                                                                                                                                                                                                                             Chordata;
Rodentia;
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                                                                                                                                 tissue-specific alternative expression in rat UDP-glucuronosyltransferase fam.
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3 POTENTIAL.
2504 MW; A221BAA6185231A4
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           abe K., Kashiwamata S.;

ng of rat liver bilirubin UDP-
cDNA: possible alternate spli
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Pred. No. 17;
2; Mismatches
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           splicing
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Murinae; Rat
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Kelley J.M.,
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I.S.,
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Best Local S
Matches 6
                                                                                                                                                                                                                                                                   01-OCT-1996
01-OCT-1996
16-OCT-2001
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CARBOHYD
CARBOHYD
                                                             Genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  use by non-profit institutions as long as its content modified and this statement is not removed. Usage by ar entities requires a license agreement (See http://www.isb or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Biochem. Biophys. Res. Commun. 169:260-264(1990).
-:- FUNCTION: UDPGT IS OF MAJOR IMPORTANCE IN THE CONJUGATION
  secretion of beta-lactamase by phosphorylation Genes Dev. 9:972-983(1995).
                                                                                                 Udo H., Munoz-Dorado J., Inouye M., Inouye S.; "Myxococcus xanthus, a Gram-negative bacterium, cortransmembrane protein serine/threonine kinase that
                                                                                                                                                                                                      Bacteria; Proteobacteria; delta Myxococcales; Cystobacterineae;
                                                                                                                                                                                                                                             PKN2
                                                                                                                                                                                                                                                     01-OCT-1996 (Rel. 34, Last sequence update) 16-OCT-2001 (Rel. 40, Last annotation updat Serine/threonine-protein kinase pkn2 (EC 2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL;
                                                                                                                                     MEDLINE-95293229; PubMed-7774814;
                                                                                                                                                    STRAIN-DZF1
                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                              Myxococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Transferase;
                                                                                                                                                                                        NCBI_TaxID=34;
                                                                                                                                                                                                                                                                                                          P54736;
                                                                                                                                                                                                                                                                                                                     PKN2_MYXXA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pfam; PF00201; UDPGT; 1.
PROSITE; PS00375; UDPGT; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    InterPro;
                                                                                                                                                                                                                                                                                                                                                                                   153
                                                                                                                                                                                                                                                                                                                                                                                                         5 PCGAVMILY 13
          IN RESPONSE TO AN EXTERNAL SIGNAL YET TO BE IDENTIFII SIMILARITY: BELONGS TO THE SER/THR FAMILY OF PROTEIN SIMILARITY: CONTAINS 1 GUANYLATE CYCLASE DOMAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           UDDGT ISOZYMES WHICH HAVE A DIFFERENT N-TERMINAL DOMAIN AN COMMON C-TERMINAL DOMAIN OF 245 RESIDUES.
SIMILARITY: BELONGS TO THE UDD-GLYCOSYLTRANSFERASE FAMILY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUBSEQUENT ELIMINATION OF POTENTIALLY ENDOGENOUS COMPOUNDS.
CATALYTIC ACTIVITY: UDP-glucuronate +
                                               FUNCTION: REGULATES THE ACTIVITY OF ENDOGENOUS BETA-LACTAMASE OF RELATED ENZYMES, BY BLOCKING THEIR SECRETION BY PHOSPHORYLATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBCELLULAR LOCATION:
ALTERNATIVE PRODUCTS:
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                                                                                                                                                                                                                                                                                                                                                                                   PCGAVLAMY 161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 D38067; BAA07262.1; -
M34007; AAA42312.1; A
Pro; IPR002213; UDPGT.
                                                                                                                                                                                                                                                                                                                                                                                                                                 6; Conserv
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531 AA;
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THE UGT1 GENE
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N-LINKED ()
N-LINKED ()
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ECB0994C6A2C7CEF CRC64;
                                                                                                                                                                                                      subdivision; Myxobacteria;
Myxococcaceae; Myxococcus.
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(GLCNAC...)
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(POTENTIAL).
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Eukaryota; Metazoa; Cl
Mammalia; Eutheria; Rc
NCBI_TaxID=10090;
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between the Swiss Institute of Bioinforth the European Bioinformatics Institute use by non-profit institutions as 1c modified and this statement is not remo
                                                                                                                                                                                                                                                                                                                                                                                                         STRAIN=NIH Swiss; TISSUE-Heart; MEDLINE-20499358; PubMed-11042143;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PROSITE; PS50125; GUANYLATE_CYCLASES_2; 1
PROSITE; PS00107; PROTEIN_KINASE_ATP; 1.
PROSITE; PS50011; PROTEIN_KINASE_DOM; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                    This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation -
                                                                                                                                                                                                                                          Genome Res. 10:1435-1462(2000).
                                                                                                                                                                                                                                                                                                        Peters T., Dildrop R., Ausmeier K., Ruether U.;
"Organization of mouse Iroquois homeobox genes in two clusters
suggests a conserved regulation and function in vertebrate
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE
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                                                                                                                                                                                                                                                                              development.
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InterPro; IPR002290; Ser_thr_pk
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                                                                                                                                                                                   SUBCELLULAR LOCATION: Nuclear (By similarity).
SIMILARITY: BELONGS TO THE TALE/IRO HOMEOBOX FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TAEPCGAITL 54
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5; Conser
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Rodentia;
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PROTEIN KINASE.
GUANYLATE CYCLASE.
ATP (BY SIMILARITY).
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Sciurognathi;
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(See http://www.isb-sib.ch/announce/
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thi; Muridae; Murinae; Mus
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                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its
                                                                                                                                                                                                                                             Peterson S., Reich C.I., McNeil L.K., Badger J.H., Glodek A., 21 Overbeek R., Goodyne J.D., Weidman J.F., McDonald L., Utterback Cotton M.D., Spriggs T., Artiach P., Kaine B.P., Sykes S.M., Sadow P.W., D'Andrea K.P., Bowman C., Fujil C., Garland S.A., Mason T.M., Olsen G.J., Fraser C.M., Smith H.O., Woese C.R.,
                                                                                                                                                                                                                                                                                                     Klenk H.-P., Clayton R.A., Tomb J.-F., White O., Nelson K.E., Ketchum K.A., Dodson R.J., Gwinn M., Hickey E.K., Peterson J.D., Richardson D.L., Kerlavage A.R., Graham D.E., Kyrpides N.C., Fleischmann R.D., Quackenbush J., Lee N.H., Sutton G.G., Gill S. Kirkness E.F., Dougherty B.A., McKenney K., Adams M.D., Loftus B. Peterson S., Reich C.I., McNeil L.K., Badger J.H., Glodek A., Zh.
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or send a
                               EMBL; AE000943; AAB88920.1; TIGR; AF2332; -.
                                                                                                                                                                                         "The complete genome sequence of the hyperthermophilic, sulphate-reducing archaeon Archaeoglobus fulgidus.";
Nature 390:364-370(1997).
                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
STRAIN-VC-16 / DSM 4304 / ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SMART; SM00389; HOX; 1.
SMART; SM00548; IRO; 1.
   Hypothetical protein; Complete SEQUENCE 148 AA; 17384 MW;
                                                                                                    use by non-profit institutions as long as its content modified and this statement is not removed. Usage by ar
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hypothetical AF2332.
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DNA_BIND 143 205 HOMEOBOX.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROSITE; PS00027; HOMEOBOX_1; PROSITE; PS50071; HOMEOBOX_2;
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                                                                                                                                                                                                                                    Venter J.C.
                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE+98049343; PubMed=9389475;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Archaeoglobaceae; Archaeoglobus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Archaea; Euryarchaeota; Archaeoglobi; Archaeoglobales
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Archaeoglobus fulgidus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; P40424; 1B7; MGI:1927642;
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9; Conserv
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IPR001356; Homeobox.
                                                                        equires a license agreement (S email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (Rel. 40, Created)
(Rel. 40, Last sequence up
(Rel. 40, Last annotation
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proteome.
8F7FF376CF10669F CRC64;
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Q9CDM7;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -i- FUNCTION: Probably phosphorylates undecaprenol to undecaprenyl
phosphate. Confers resistance to bacitracin (By similarity).
-i- CATALYTIC ACTIVITY: ATP + undecaprenol - ADP + undecaprenyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               lactis ssp. lactis IL1403.";
Genome Res. 11:731-753(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=21235186; PubMed=11337471;
Bolotin A., Wincker P., Mauger S., Jaillon O., Malarme K.,
Weissenbach J., Ehrlich S.D., Sorokin A.;
"The complete genome sequence of the lactic acid bacterium Lactococcus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ÜPK OR BACA OR LL2192.
Lactococcus lactis (subsp. lactis) (Streptococcus lactis).
Bacteria; Firmicutes; Lactobacillales; Streptococcaceae; Lactococcus.
NCBI_TaxID=1360;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         15-JUN-2002 (Rel. 41, Greated)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Putative undecaprenol kinase (EC 2.7.1.66) (Bacitracin resistance)
                                                                                                                                                                                                                                                                                                                                                                                                                                Transferase; Kinase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pfam;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; AE006448; AAK06290.1; -.
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    57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phosphate.
SUBCELLAULAR LOCATION: Integral membrane protein (Probable)
MISCELLANEOUS: Bacitracin is thought to be involved in inh
of peptidoglycan synthesis by sequestering undecaprenyl
diphosphate reducing the pool of lipid carrier available.
SIMILARITY: BELONGS TO THE UPK FAMILY.
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    SVVVLYFHK
                                                AVMILYFHK 16
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                                                                                            6; Conserv
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7; Conser
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2284 AA;
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                                                                                               Conservative
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                                                                                               ω,
                                                                                                                                                                                                                                                             POTENTIAL.
POTENTIAL.
POTENTIAL.
POTENTIAL.
                                                                                               Pred. No. 26;
3; Mismatches
                                                                                                                    Score 39; DB Pred. No. 26;
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83ABD014DDDBB14E CRC64;
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01-AUG-1992 (Rel. 2
01-NOV-1995 (Rel. 3
15-JUL-1998 (Rel. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
MEDLINE=92124734; Pubb
Renne R., Friedl E., 9
Neumann-Haefelin D.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              15-DEC-1998
15-DEC-1998
15-JUN-2002
MEDLINE-98044033; PubMed-9384377;
Kunst F. Ogasawara N., Moszer I., Albertini A.M., Alloni G.,
Azevedo V., Bertero M.G., Bessieres P., Bolotin A., Borchert S.,
Borriss R., Boursier L., Brans A., Braun M., Brignell S.C., Bron S.,
Brouillet S., Bruschi C.V., Caldwell B., Capuano V., Carter N.M.,
Choi S.K., Codani J.J., Connerton I.F., Cummings N.J., Daniel R.A.,
Denizot F., Devine K.M., Dusterhoft A., Ehrlich S.D., Emmerson P.T.
Entian K.D., Errington J., Fabret C., Ferrari E., Foulger D.,
Fritz C., Fujita M., Fujita Y., Fuma S., Galizzi A., Galleron N.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EMBL; M74895; AAA47800.1; -.
EMBL; M74895; AAA47797.1; ALT_INIT.
PIR: E40820; WMLJBT.
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Hypothetical protein.
SEQUENCE 388 AA; 44527 MW
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Viruses; Retroid vi
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                                                                                                                                                                                                                                                                                                                        STRAIN-168;
                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A
                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Firmicutes; NCBI_TaxID=1423;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacillus subtilis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 precursor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypothetical
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1 ABC transporter extracel)
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RESULT 15
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or send a
                                                                              KIN2_YEAST STANDARD; PRT; 1147 AA P13186; Q12384; 01-JAN-1990 (Rel. 13, Created) 15-DEC-1998 (Rel. 37, Last sequence update) 30-MAY-2000 (Rel. 39, Last annotation update) Protein kinase KIN2 (EC 2.7.1.-). KIN2 OR YLR096W OR L8004.3.
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Nature 390:249-256(1997).
-!- FUNCTION: PROBABLY PART OF A BINDING-PROTEIN-DEPENDENT TRANSPORT
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SubtiList; BG14001; yuro.
InterPro; IPR000567; SBP_bac_1.
Pfam; PF01547; SBP_bacterial_1; 1.
PROSITE; PS00013; PROKAR_LIPOPROTEIN; 1.
PROSITE; PS01037; SBP_BACTERIAL_1; FALSE_NEG_PROSITE; PS01037; SBP_BACTERIAL_1; FALSE_NEG_Hypothetical protein; Transport; Membrane; Li
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                       Saccharomyces cerevisiae (Baker's yeast).
Eukaryota: Fungi: Ascomycota: Saccharomycotina: Saccharomycetes:
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an email to license@isb-sib.ch).
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Saccharomyces
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Wilson R., Waterston R.;
Submitted (APR-1996) to the EMBL/GenBank/DDBJ databases.
Submitted (APR-1996) THE PROTEIN IS PROBABLY A SERINE/THREONINE
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EMBL; Z73269; CAA97661.1; -.
EMBL; U53876; AAB67540.1; -.
EMBL; U53876; AAB67540.1; -.
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SMART; SM00220; S_TKC; 1.
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InterPro; IPR001772; Kinase_Cterm.
InterPro; IPR002290; Ser_thr_pkinase.
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Q96033; O1-DEC-2001 (TrEMBLrel. 19, Created)
O1-DEC-2001 (TrEMBLrel. 19, Last sequence update)
O1-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Hypothetical 17.5 kDa protein.
B208.030.
                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.

Schulte U., Aign V., Hoheisel J., Brandt P., Fartmann B.

Nyakatura G., Mewes H.W., Mannhaupt G.;

Submitted (MAY-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                       Neurospora crassa.
Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycetes;
Sordariales; Sordariaceae; Neurospora.
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Submitted (SEP-2001) to the Ex
EMBL; ACO93703; AALOO863.1; -
InterPro; IPR001980; SH2.
InterPro; IPR001452; SH3.
Pfam; PF00017; SH2; 1.
Pfam; PF00018; SH3; 1.
PROSITE; PS50002; SH3; 1.
                                                                                                               SÉDUENCE FROM N.A.

STRAIN-AV19 / DSM 6324 / JCM 9639;

MEDLINE-21927647; PubMed-11930014;

Slesarev A.I., Mezhevaya K.V., Makarova K.S., Polushin N.N.,

Shcherbinina O.V., Shakhova V.V., Belova G.I., Aravind L.,

Natale D.A., Rogozin I.B., Tatusov R.L., Wolf Y.I., Stetter K.O.,

Malykh A.G., Koonin E.V., Kozyavkin S.A.;

"The complete genome of hyperthermophile Methanopyrus kandleri AV19

and monophyly of archaeal methanogens.";

Proc. Natl. Acad. Sci. U.S.A. 99:4644-4649(2002).

EMBL: AE010365; AAM01936.1; -.
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Rhabditidae; Peloderinae;
NCBI_TaxID=6239;
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01-JUN-2002 (TrembLrel.
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Archaea; Euryarchaeota;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Science
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Q9V2Z9;
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Archaea; Euryarchaeota; Archaeoglobi;
Archaeoglobaceae; Archaeoglobus.
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01-JAN-1998 (TrEMBLrel. 05, 1
01-JUN-2002 (TrEMBLrel. 21, 1
Signal-transducing histidine
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Pfam; PF02249; MCR_alpha; 1.
Pfam; PF02745; MCR_alpha_N; 1.
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FEMS Microbiol Lett 177:101-108(1999).
EMBL; AFI21100; AA045632.1; -.
HSSP; P11558; 1MRO.
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"A phylogenetic analysis of microbial conmethane hydrate containing marine fluids
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Magnaporthe grisea (Rice blast fungus) (Pyricularia grisea). Eukaryota; Fungi; Ascomycota; Pezizomycotina; Sordariomycete Sordariomycetes incertae sedis; Magnaporthaceae; Magnaporthe NCBI_TaxID-148305;
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Nature 390.364-370(1997).
EMBL; AE001001; AAB89779.1; -.
TIGR; AF1467; -.
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SMART; SM00388; HiskA; 1.
Hypothetical protein; Complete proteome.
SEQUENCE 222 AA; 25036 MW; 1B7B4E1B9FB70705 CRC64;
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InterPro; IPR003661; His_kinA.
InterPro; IPR004359; HIS_KIN_sig.
Pfam; PF02518; HATPase_C; 1.
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MEDLINE-21886394; PubMed-11889109;
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J. Bacteriol. 184:2005-2018(2002).
EMBL; AE010543; AAL94512.1; -.
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01-MAR-2001 (TrEMBLrel. 16, Created)
01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation updat
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SEQUENCE FROM N.A.
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STRAIN-INA168; TRANSPOSON-OCCAN;
Kito H., Takahashi Y., Sato J., Sone T., Tomita F.;
"Occan, a DNA-type transposon sequence identified fi
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Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databi
EMBL; AB074754; BAB784491.; -
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The Sequence of the Ovine Gamma-Carboxylase cDNA.
Submitted (OCT-2000) to the EMBL/GenBank/DDBJ data
EMBL; AF312035; AAG30935.1; -.
InterPro; IPR001870; Gamma-Carbxylse.
InterPro; IPR000504; RNA_rec_mot.
PROSITE; PS00030; RRW_RNP_1; UNKNOWN_1.
SEQUENCE 758 AA; 87581 MW; 87F8B58C187B9091 CR
SEQUENCE FROM N.A.

MEDILINE=21948211; PubMed=11950982;

Springer N.M., Danilevskaya O.N., Hermon P., Helentjaris T.

Springer N.M., Campler H.F., Kaeppler S.M.;

"Sequence Relationships, Conserved Domains, and Expression for Maize Homologs of the Polycomb Group Genes E(z), esc, a
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Eukaryota; Viridiplantae;
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MEDLING-21595285; PubMed-11759840;

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Watanabe A., Iriguchi M., Ishikawa A., Kawashima K., Kimura

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Kishida Y., Shimpo S., Sugimoto M., Takazawa M., Yamada M.,
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01-JUN-2002 (TrEMBLrel. 21, Last sequence up
01-JUN-2002 (TrEMBLrel. 21, Last annotation
Sphingolipid ceramide N-deacylase.
Shewanella algae.
                                                            Deacylase from a Marine Bacterium,
J. Biol. Chem. 0:0-0(2002).
EMBL; AB079849; BAB85110.1; -.
SEQUENCE 992 AA; 109848 MW; CEK
                                                                                                                                                               Furusato M., Sueyoshi N., Mitsutake S., Sakaguchi K., Ki
Okino N., Ichinose S., Omori A., Ito M.;
"Molecular Cloning and Characterization of Sphingolipid
Deacylase from a Marine Bacterium, Shewanella alga G8.";
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DNA Res. 8:205-213(2001).
EMBL; AP003594; BAB75469.1;
Hypothetical protein; Comple
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01-MAR-2002
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EMBL; AF443597; AAM13421.1; -.
                                                                                                                                                                                                                                                                                                         STRAIN-G8;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Shewanella.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Proteobacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QBRTY6;
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Bacteria; Cyanobacteria; Nostocales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hypothetical protein ASR3770.
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                  Fraser C.M., Norris S.J., Weinstock G.M., White O., Sutton G.G. Dodson R., Gwinn M., Hickey E.K., Clayton R., Ketchum K.A., Sodergren E., Hardham J.M., McLeod M.P., Salzberg S., Peterson Khalak H., Richardson D., Howell J.K., Chidambaram M., Utterbac McDonald L., Artiach P., Bowman C., Cotton M.D., Fujii C., Garl Hatch B., Horst K., Roberts K., Sandusky M., Weidman J., Smith
                                                                                                                                                               EMBL;
HSSP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sueyoshi N., Furusato M., Okino N., Mitsutake S., Ito M.; "Molecular cloning and characterization of sphingolipid c deacylase from a marine bacterium, Shewanella alga NS589. Submitted (FEB-2002) to the EMBL/GenBank/DDBJ databases. EMBL; AB079850; BAB85111.1; -. 87611F6BE9DAEBBA CRC64;
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MEDLINE=98332770; PubMed=9665876;
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Bacteria; Spirochaetales;
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                                                           PF00580; UvrD-helicase;
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8; Conserv
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2001259; AAC65853.1;
   proteome.
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MEDLINE-20150255; PubMed-10684935;
MEDLINE-20150255; PubMed-10684935;
Mhite O., Hickey E.K., Peterson J., Utterback T., Berry
Linher K., Weidman J., Khouri H., Craven B., Bowman C.,
Gwinn M., Nelson W., DeBoy R., Kolonay J., McClarty G.,
"Genome sequences of Chlamydia trachomatis MoPn and Chla
                                                                                                                          SET domain protein.
CPJ0878 OR CP0991.
Chlamydia pneumoniae (Chlamydophila pneumoniae).
Bacteria; Chlamydiales; Chlamydiaceae; Chlamydophila.
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SEQUENCE
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STRAIN-K-12: TRANSPOSON-TN7;

Sampei G., Motomura K., Masuda S., Yamaguchi T.

Furuya N., Komano T., Mizobuchi K.;

Furuya N., Komano T., Mizobuchi K.;

"Organization and diversification of plasmid g

nucleotide sequence of the R721 genome.";

submitted (JUN-2000) to the EMBL/GenBank/DDBJ
                                                                                                                                                                        01-OCT-2000 (TREMBLIGEL 15, 01-OCT-2000 (TREMBLIGEL 15, 01-OCT-2001 (TREMBLIGEL 18,
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STRAIN-K-12; TRANSPOSON-TN7;
MEDLINE-93015772; PubMed-1400257;
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01-MAR-2001 (TrEMBLrel.
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Plasmid R721.
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                                                                                   STRAIN-AR39;
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                                                                                                                                                                                                                                                                                                                                                                                                    Nucleotide sequence of the R721 shufflon."; Bacteriol. 174:7053-7058(1992).
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Search completed: November Job time : 31 secs

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                                 Query Match
Best Local Similarity
Matches 6; Conserv
                                                                                                         "Comparison of whole genome sequences of Chlamydia pneumoniae J138 from Japan and CWLO29 from USA.";
Nucleic Acids Res. 28:2311-2314(2000).
EMBL; AE002257; AAF38770.1;
EMBL; AP002548; BAA99086.1;
InterPro.
                                                                     Pfam; PF00856; SET; 1.
SMART; SM00317; SET; 1.
PROSITE; PS50280; SET; 1.
SEQUENCE 221 AA; 25889
                                                                                                                                                                                                                 pneumoniae AR39.*;
Nucleic Acids Res.
[2]
                                                                                                                                                                                                         SEQUENCE FROM N.A.
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TTEPCSSIHISLNNDWR
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